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Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: [year=2010; month=8; day=10; hr=16; min=0; sec=33; ms=891;]

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Application No: 10590936 Version No: 1.0

Input Set:

Output Set:

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Finished: 2010-08-05 11:54:00.915
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 468 ms
Total Warnings: 4
Total Errors: 0
No. of SeqIDs Defined: 8
Actual SeqID Count: 8

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SEQUENCE LISTING

<110> TECNOGEN S.C.P.A.
DE SANTIS, Rita
PALOMBO, Giovanna
CARMINATI, Paolo
PELLICCIA, Angela

<120> ANTI-HUMAN TENASCIN MONOCLONAL ANTIBODY

<130> 725.1051

<140> 10590936

<141> 2010-08-05

<150> PCT/IT2005/000078

<151> 2005-02-16

<150> RM2004A0000105

<151> 2004-02-27

<160> 8

<170> PatentIn version 3.5

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<221> CDS

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gtc ata atg tcc aga gga caa att gtt ctc tcc cag tct cca gca atc	96
Val Ile Met Ser Arg Gly Gln Ile Val Leu Ser Gln Ser Pro Ala Ile	
20 25 30	

ctg tct gca tct cca ggg gag aag gtc aca atg act tgc agg gcc aac	144
Leu Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Arg Ala Asn	
35 40 45	

tca agt gta cgt ttc atg cac tgg tac cag cag aag cca gga tcc tcc	192
Ser Ser Val Arg Phe Met His Trp Tyr Gln Gln Lys Pro Gly Ser Ser	
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ccc aaa ccc tgg att tat gcc aca tcc aac ctg gct tct gga gtc cct	240
Pro Lys Pro Trp Ile Tyr Ala Thr Ser Asn Leu Ala Ser Gly Val Pro	
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gct cgc ttc agt ggc agt ggg tct ggg acc tct tat tct gtc aca atc	288
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Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Val Thr Ile
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agc aga gtg gag gct gaa gat gct gcc act tat tac tgc cag cag tgg 336
Ser Arg Val Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp
100 105 110

agt agt aat tca ccc agg acg ttc ggt gga ggc acc aag gtg gaa atc 384
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Leu Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Arg Ala Asn
35 40 45

Ser Ser Val Arg Phe Met His Trp Tyr Gln Gln Lys Pro Gly Ser Ser
50 55 60

Pro Lys Pro Trp Ile Tyr Ala Thr Ser Asn Leu Ala Ser Gly Val Pro
65 70 75 80

Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Val Thr Ile
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Arg Arg Ala
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 gtc cac tct gag gtc cag ctg caa cag tct gga cct gag ctg gtg aag 96
 Val His Ser Glu Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys
 20 25 30
 cct gga gct tca atg aag att tcc tgc aag gct tct ggt tac tca ttc 144
 Pro Gly Ala Ser Met Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe
 35 40 45
 act ggc tac acc atg aac tgg gtg aag cag agc cat gga aag aac ctt 192
 Thr Gly Tyr Thr Met Asn Trp Val Lys Gln Ser His Gly Lys Asn Leu
 50 55 60
 gaa tgg att gga ctt att aat cct cac aat ggt ggt act acc tac aac 240
 Glu Trp Ile Gly Leu Ile Asn Pro His Asn Gly Gly Thr Thr Tyr Asn
 65 70 75 80
 cag aag ttc aag ggc aag gcc aca tta act gta gac aag tca tcc aac 288
 Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Asn
 85 90 95
 aca gcc tac atg gag ctc ctc agt ctg aca tct gag gac tct gca gtc 336
 Thr Ala Tyr Met Glu Leu Leu Ser Leu Thr Ser Glu Asp Ser Ala Val
 100 105 110
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 Tyr Tyr Cys Thr Arg Pro Gly Gly Tyr Tyr Trp Phe Phe Asp Val Trp
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Pro Gly Ala Ser Met Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe
35 40 45

Thr Gly Tyr Thr Met Asn Trp Val Lys Gln Ser His Gly Lys Asn Leu
50 55 60

Glu Trp Ile Gly Leu Ile Asn Pro His Asn Gly Gly Thr Thr Tyr Asn
65 70 75 80

Gln Lys Phe Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Asn
85 90 95

Thr Ala Tyr Met Glu Leu Leu Ser Leu Thr Ser Glu Asp Ser Ala Val
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<212> DNA

<213> Artificial Sequence

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